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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/998,350

DATE: 12/23/2002 P.6

TIME: 14:13:23

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\12232002\I998350.raw

```
3 <110: APPLICANT: Roller, Peter P
             Long, Ya-liu
     1.
             Lung, Feng-Di T
             Hing, Elenter C
             Mang, Dajun
     ** <12** TITLE OF INVENTION: PEDCX-STABLE, NON-PHOSPHORYLATED CYCLIC PEPTIDE INHIBITORS
OF SH2 DOMAIL
            BINDING TO TARGET PROTEIN, CONJUGATES THEREOF, COMPOSITIONS AND METHODS OF
     SYNTHESIS AND USE
     15 () to FILE REFERENCE: 1146-3
     15 CLASS TURRENT APPLICATION NUMBER: 09/998,350
C--> 16 <141> CURRENT FILING DATE: 2002-12-09
     18 CHAR PEACE APPLICATION NUMBER: PCT/US00/15201
     18 KIND PRICE BULING DATE: 2000-06-02
     20 GLADA PERCE APPLICATION NUMBER: 60/137,187
     24 (1000 MINERE OF SEQ ID NOS: 19
     20 COURSETWARE: Patentin version 3.1
    28 KH 0 SEQ ID NO: 1
     29 KEIDS LENGTH: 9
     30 CHLO CYPE: ERT
     39 K.H. & OFGANISM: Artificial Sequence
     BEATTER:
     34 K.E. S. OTHER INFORMATION: Synthetic
     36 KILLS SEATURE:
     3" Kill Do MAME FFY: musc feature
     38 (M. D. ICCATION: 11)...(1)
     30 < 1 0 OTHER INFORMATION: Maa = Gla, which is gamma-carboxy-L-glutamic acid
     42 (LIO EFATURE:
     49 Kin D NAME REY: masc feature
     44 (513) LCCATION: (4)..(9)
     45 Chr. CIHER INFIEMATION: Tyr at position 9 is an amide, i.e. C(O)NH
     48 Chich FEATURE:
     49 KM D NAME KHY: musc feature
     50 < 0.10 \text{ IOCATION:} 1)..(9)
     51 KALUS OTHER INFORMATION: Maa (Gla) and Tyr at position 9 are bridged together, making
this
             peptide cyclic
     55 < 40.00 SEQUENCE: 1
W--> 57 Xaa Leu Tyr Glu Asn Val Gly Met Tyr
    5h \cdot 1
     61 <210: SEQ ID NO: 2
     62: <2.1: LENGTH: 9
```

63 <212/ TYPE: PRT



64 <213> ORGANISM: Artificial Sequence

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```
Input Set : A:\PTO.VSK.txt
                      Output Set: N:\CRF4\12232002\I998350.raw
     66 <2005 FEATURE:
     67 <2.7. OTHER INFORMATION: Synthetic
     69 (3.1) FEATURE:
     70 CL E MAME MEY: mass feature
     71 (d. . / DOMATION: (1:..(1)
      The CLAR OLEGE INFORMATION: Mag at position 1 is alpha-amino-adipic acid (Adi)
     95 Ch : PEATTEE:
     74 Ct. 1.8 DAME TEY: most feature
     77 (1) 36 MITON: (4)...(4)
     78 CL + OTHER INFORMATION: Xaa at position 4 is Glu or Adi
     EL COUR FEATURE:
     E. C. L. MAME HEY: masc feature
     \epsilon s of the property of \epsilon . (9)
     E4 CLTS OTHER INFORMATION: Tyr at position 9 is an amide, i.e., C(O)NH ET CLTS PEATURE:
     \xi_{\text{T}} \ll 1.1 + \text{MAME/FFY:} \pi_{\text{BBG}} feature
     89 (22) - LO MATRON: (11...(9)
     90~\odot...~0THER INFORMATION: Mas at position 1 and Tyr at position 9 are bridged together,
mak
              ing this peptide cyclic
     91 (1 ). SEQUENCE: 2
W--> 96 Xaa Leu Tyr Xaa Asn Val Gly Met Tyr
     9.1
     100 K. D. J. SHOW IF NO: 3
     101 (L.1) IENGTH: 9
     1 C. C.L. PYPE: PRI
     107 table ON WANTEM: Artificial Sequence
     100 OLLOS FRATURE:
     100 C. O DUBER INFORMATION: Synthetic
     100 C.O. FWATUFF:
     1 0 K. D. MAME, PRY: musc feature
     110 d.10 DOCATION: (1)...(1)
     111 KINDS OTHER INFORMATION: Kaa is any amino acid other than Glu
     114 C'US FEATURE:
     115 K. D. NAME/EFY: musc feature
     in the thermal (0.01, 0.03)
     1:0 K ADD OTHER INFORMATION: Tyr at position 9 is an amide, i.e., C(O)NH
     1.0 < .100 EFATURE:
     111 C. H. MAME, FEY: musc feature
     111 GREE LOCATION: (1)...(9)
     11.2 CLEEN CHEEK INFORMATION: Xaa and Tyr at position 9 are bridged together, making this
pepti
              - de cyclic
     10 <4000 SEQUENCE: 3
W--> 129 Xaa Leu Tyr Glu Asn Val Gly Met Tyr
     1 > 1
                          5
     133 <1400 SEC II NO: 4
     134 G 110 LENGTE: 10
     101 G 120 TYPE: PAT
     196 CLUB CRGANISM: Artificial Sequence
     158 < 20> FEATURE:
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/998,350

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TIME: 14:13:23

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Input Set : A:\PTO.VSK.txt
                      Gutrut Set: N:\CRF4\12232002\I998350.raw
     149 <2205 OTHER INFORMATION: Synthetic
     141 <2200 FEATURE:
     142 KORD MAME/FEY: musc_feature
     14 ' <1711 IOCATION: (1)...(1)
     144 KINES OTHER INFOFMATION: Xaa = Gla, which is gamma-carboxy-L-glutamic acid
     14. C.O. BEATURE:
     14% K.IID MAME/FEY: masc feature
     14 · <0000 LOCATION: (10)...10)
     19. C. TO OTHER INFORMATION: Cys at position 10 is an amide, i.e., C(O)NH
     1' COURT FEATURE:
     194 C. ... MAME/FEY: musc feature
     1.6 disconfidential: (1)..(10)
     150 C. TO OTHER INFORMATION: Xaa (Gla) and Cys are bridged together, making this peptide
cycli
     1:
     160 K4 GD SEQUENCE: 4
W--> 162 Xaa Leu Tyr Glu Asn Val Gly Met Tyr Cys
     166 K. 100 SEC 01 NO: 5
     1... <. 11: DENGTH: 19
     168 K. L. TYLE: ERT
     169 K. 180 ORFANISM: Artificial Sequence
     171 K. DE PEATURE:
15 COST OTHER INFORMATION: Symthetic
     1/4 C. : PEATURE:
     1 * C. D. MASTE/KEY: musc feature
     116 < ...  100 \text{ ATION: } (1)...(1)
     10^{\circ} K. \odot OTHER INFORMATION: Xaa = Gla, which is gamma-carboxy-L-glutamic acid
     100 K4: 1 FEGUENCE: 5
W--> 182 Xaa Leu Tyr Glu Asn Val Gly Met Tyr Cys
     1 · · · l
     186 C. 103 J.D. II NO: 6
     18 For Lib LENGIH: 11
     1 mm of the CYPE: FRT
     1- + KAR OF (OS) AND SM: Artificial Sequence
     141 1.200 PEACUEE:
     1. CAMP OTHER INFORMATION: Synthetic
     1 44 0.000 FEATURE:
     1 to 4.200 MAME, FER: musc feature
     1 0 < 0.000 LOCATION: (1)...(1)
     190 CLMS CLMEE INFOFMATION: Xaa = Gla(OtBu)2, which is di-tert-butoxy-gamma-carboxy-L-
glutam
     1.4:
                ur acid
     201 GRO FEATURE:
     20% KLBD MAME, MEY: misc feature
     205 < 200 \text{ IOCATION: } (5)...(3)
     20; CLES OTHER INFOFMATION: Tyr at position 3 is modified to Tyr(tBu), which is tert-
but.yl-**
     200
               romine
     208 < 200 FEATURE:
     209 <1210 NAME, FEY: misc feature
     210 < .22> LOCATION: (4)..(4)
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/998,350

DATE: 12/23/2002

TIME: 14:13:23

```
Input Set : A:\PTO.VSK.txt
                                              Cutput Set: N:\CRF4\12232002\I998350.raw
           211 <2235 FINER INFORMATION: Glu at position 4 is modified to Glu(OtBu), which is tert-
butox/-
          .:12
                                edutamic acid
           .15 < 200 FEATURE:
           .:16 < 21: MAME FEY: namec_feature</pre>
           217 < 0.00 < 100 ATION: -5)...(5)
          113 C. A FIEER INFOFMATION: Asn at position 5 is modified to Asn(Trt), which is is
trytyl-usp
          .:1:4
                                 .:i :g:r.∈
          .12. C. O. FEATURE:
           May as the MAMERET: masc feature
           Und C. 10 LOWATION: (9)...(9)
           Mark Court of HER INFORMATION: Tyr at position 9 is modified to Tyr(tBu), which is tert-
butyl-*y
          ...
                                ::.in∈
           H: C.O. PEATURE:
           do: C.D. DAME TEY: raise feature
           that we in LOCATION: (10) \dots (10)
          AND RELATED INFOFMATION: Cys at position 10 is modified to Cys(Trt), which is trytyl-
cyste
                                 in, and Tys(Trt) is connected to a resin
           Associated ABOUFNEE: 6
W--> 238 Xaa Leu Tyr Glu Asn Val Gly Met Tyr Cys
           1.50
           Man Color OFR II NO: "
           34 KUNIN LEMGTE: 9
           344 C. III TYPE: PRT
           34 Clo +8 AMISM: Artificial Sequence
           Barris of BEATTEE:
           142 K. S . THER INFERNATION: Synthetic
           DO COUR FEATURE:
           uni d. I MAME MEY: misc feature
           (ii), C_{1} > C_{2} > C_{3} > C_{4} 
            186 C. C. THER INFORMATION: Xaa = Gla, which is gamma-carboxy-L-glutamic acid
           MO CLUD HEATHER
           AND KILLS DAME REY: risc feature
           MORE COLD FOR ACTION: (1)...(1)
           .5 * <...: ** THEE INFORMATION: Xaa has a ClCH2C(O) - group attached</p>
            ON COUNTY SEATURE:
           . Who call NAME 'KEY: masc feature
          1.64 \times 1.12 \times 10^{\circ} \text{ATION:} \text{ (9)}
           1800 C.E. CIHER INFORMATION: Tyriat position 9 has a -C(CH2SH)C(O)NH2 group attached
           THE CHOICE SEQUENCE: 7
W--> 270 Xaa Leu Tyr Glu Asn Val Gly Met Tyr
             74 0.100 PFG HENO: 8
                  c. 110 LENGTE: 10
             1.0 < UD TYPE: PHT
             V < 100 + EGANISM: Artificial Sequence
             1 + <.20> :EATURE:
           380 K. 250 CTHER INFORMATION: Synthetic
           382 <120> FEATURE:
           383 < 21> NAME, KEY: misc feature
```

RAW SEQUENCE LISTING

FATENT APPLICATION: US/09/998,350

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DATE: 12/23/2002
                     RAW SEQUENCE LISTING
                     PATENT APPLICATION: US/09/998,350
                                                            TIME: 14:13:23
                     Input Set : A:\PTO.VSK.txt
                     Cutput Set: N:\CRF4\12232002\I998350.raw
     284 <222> LOCATION: -1 ... (1)
     2:5 <223 CTHEE INFOFMATION: Xaa - Adi, which is alpha-amino-adipic acid
     250 CRRO FEATURE:
     200 C2210 NAME MEY: rubb_feature
     100 CMM DOCATION: (1 ...(1)
     1941 0.25
               THEE INFOFMATION: Xaa has a CH2CO- group attached
     194 COSTO PEATMEE:
     190 COMM NAME/HEY: rusc_feature
     1900 CARL TOTALION: (10)...101
     190 C.B. THEE INFOFMATION: Cys is an amide, i.e., C(0)NH
     POST COMPARES
     NOT 4 HILL MAME REY: raid feature
     96. CHIC TOCATION: (10..(10)
     Post CLASS OTHER INFORMATION: Maa (Adi) and Cys are bridged together, making this peptide
cycli
     1 .
     1.00
        - GOUL L'EQUENCE: 8
W--> 309 Xaa Leu Tyr Glu Asn Val Gly Met Tyr Cys
     10.10
     Tire Click . E. H. NO: 9
     014 COLD LENGTH: 10
     BIT COLD TYPE: FET
     Plo Krist FSAMISM: Amificial Sequence
     PIL COMMUNICATIVEF:
     Fig. ( )) - THEF INFOFMATION: Synthetic
     N. . CHOO FEATURE:
     b.. 0.000 NAME/FFY: raid_feature
     0, + 0.00 \in 100 ATION: (1)...(1)
     E.4 CLEAR CHEEK INFORMATION: At position 1, Xaa = Adi, which is alpha-amino-adipic acid
     ROUGH STREET
     PDF 0.000 MAME/FEY: rand_feature
     En • 3. 11 0 10 TATION: (4) ...(4)
     From CHILL OTHER INFORMATION: At position 4, Xaa = Adi, which is alpha-amino-adipic acid
     Barrier HOUSE FEATURE:
     Pod K. 1819 DAME FEY: musc feature
      For K.EEC CHEEL INFORMATION: Cys is an amide, i.e., C(0)NH
      S C C 201 PEATURE:
     940 K. 200 MAME HEY: masc feature
     341 < 221 \text{ LOMATION: } (1)...(10)
     241 C.231 OTHER INFORMATION: Xaa (Adi) at position 1 and Cys are bridged together, making
this
              peptide cyclic
     (4) (400) PEQUENCE: 9
W--> 348 Xaa Leu Tyr Xaa Asn Val Gly Met Tyr Cys
     249 1
     [12] <. 10: . EQ ID NO: 10
     [53 < 11] [ENGTH: 10
     [[4] <, 120 CYPE: PRT
     555 < 13: ORGANISM: Artificial Sequence
     357 < . 20> FEATURE:
```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/998,350

DATE: 12/23/2002
TIME: 14:13:24

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\12232002\I998350.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1 Seq#:2; Xaa Pos. 1,4 Seq#:3; Xaa Pos. 1 Seq#:4; Xaa Pos. 1 Seq#:5; Xaa Pos. 1 Seq#:6; Xaa Pos. 1 Seq#:7; Xaa Pos. 1 Seq#:8; Xaa Pos. 1 Seq#:9; Xaa Pos. 1,4 Seq#:10; Xaa Pos. 8 Seq#:11; Xaa Fos. 10 Seq#:12; Xaa Fos. 8 Seq#:13; Xaa Fos. 8,10 Seq#:14; Xaa Sos. 10 Seq#:18; Xaa Fos. 1 Seq#:19; Xaa Bos. 1

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 9

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/998,350

DATE: 12/23/2002 TIME: 14:13:24

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\12232002\I998350.raw

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0 I: 96 M: 341 W: (46: "n" or "Xaa" used, for SFQ 1D#:2 after pos.:0 L:129 M:341 W: (46) "n" or "Kaa" used, for SEQ 1D#:3 after pos.:0 L:162 M:341 W: (46: "n" or "Xaa" used, for SEQ ID#:4 after pos.:0 L:182 M:341 W: (46) "n" or "Kaa" used, for SEQ ID#:5 after pos.:0 L::38 M:341 W: (46: "n" or "Kaa" used, for SEQ ID#:6 after pos.:0 L::70 M:341 W: (400 "n" or "Kaa" used, for SEC ID#:7 after pcs.:0 L:500 M:341 W: (460 "n" or "Kaa" used, for SEC ID#:8 after pos.:0 L:348 M:341 W: (400 "n" or "Kaa" used, fir SEC 1D#:9 after pos.:0 L:580 M:541 W: (400 "n" or "Xaa" used, for SEQ ID#:10 after pos.:0 L:441 M:341 W: (4++ "n" or "Kaa" used, for SEC ID#:11 after pos.:0 L:474 M:541 W: (4): "n" or "Kaa" used, for SEC ID#:18 after pos.:0 L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0 L:557 M:341 W: (4) "n" or "Kaa" used, for SEQ ID#:14 after pos.:0 L:641 M:341 W: (46) "n" or "Kaa" used, for SEC ID#:18 after pos.:0 L:677 M:341 W: (40) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0